
Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: Mon Oct 22 17:32:19 EDT 2007

Reviewer Comments:

Ile Ala Gly Ser Ala Thr Val Lys Ile Gly Glu Lys Val His Glu Ile
245 250 255

Gly Ile Ala Gly Lys Gln 260

1

1

Please delete the above end of file errors at the end of sequence id#12.

Validated By CRFValidator v 1.0.3

Application No: 10574424 Version No: 1.0

Input Set:

Output Set:

Started: 2007-10-03 15:38:28.689

Finished: 2007-10-03 15:38:30.099

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 410 ms

Total Warnings: 12

Total Errors: 4

No. of SeqIDs Defined: 12

Actual SeqID Count: 12

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W	402	Undefined organism found in <213> in SEQ ID (2)								
W	402	Undefined organism found in <213> in SEQ ID (3)								
W	402	Undefined organism found in <213> in SEQ ID (4)								
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E	321	No. of Bases conflict, this line has no nucleotides SEQID (12)								
E	355	Empty lines found between the amino acid numbering and the								
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Asn Gly Phe Lys Ala Gly Glu Thr Ile Tyr Asp Ile Asp Glu Asp Gly
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                             105
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Leu Ala Asp Thr Asp Ala Ala Leu Asp Ala Thr Thr Asn Ala Leu Asn
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Asr	n Thr	Lys 195	Ala	Asp	Glu	Ala	Val 200	Lys	Thr	Ala	Asn	Glu 205	Ala	Lys	Gln
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Thr	Ala 50	Ala	Asp	Val	Glu	Ala 55	Asp	Asp	Phe	Lys	Gly 60	Leu	Gly	Leu	Lys
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Leu Leu 65	Leu Gly	Gln Val	. Ala	Thr	Glu	Gly	Glu 75	Lys	Gln	Phe	Val	Gly 80
Gln Ile	Ala Arg	Ser Glu 85	ı Gln	Ala	Ala	Glu 90	Gly	Val	Tyr	Asn	Tyr 95	Ile
Thr Val	Ala Ser 100	Leu Pro	Arg	Thr	Ala 105	Gly	Asp	Ile	Ala	Gly 110	Asp	Thr
Trp Asn	Thr Ser	Lys Val	. Arg	Ala 120	Thr	Leu	Leu	Gly	Ile 125	Ser	Pro	Ala
Thr Gln 130	Ala Arg	Val Lys	: Ile 135	Val	Thr	Tyr	Gly	Asn 140	Val	Thr	Tyr	Val
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Ser Thr	Thr Val	Gly Val	. Gln	Lys	Val	Ile 170	Thr	Leu	Tyr	Gln	Asn 175	Tyr
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Val Asp Gly Asn Leu Thr Met His Gly Lys Thr Ala Pro Val Lys Leu

Lys Ala Glu Lys Phe Asn Cys Tyr Gln Ser Pro Met Glu Lys Thr Glu 120 Val Cys Gly Gly Asp Phe Ser Thr Thr Ile Asp Arg Thr Lys Trp Gly 130 135 Met Asp Tyr Leu Val Asn Val Gly Met Thr Lys Ser Val Arg Ile Asp 155 Ile Gln Ile Glu Ala Ala Lys Gln 165 <210> <211> 464 <212> PRT <213> N. meningitidis Ser Pro Asp Val Lys Ser Ala Asp Thr Leu Ser Lys Pro Ala Ala Pro 5 10 Val Val Ser Glu Lys Glu Thr Glu Ala Lys Glu Asp Ala Pro Gln Ala 20 25 Gly Ser Gln Gly Gln Gly Ala Pro Ser Ala Gln Gly Ser Gln Asp Met 40 Ala Ala Val Ser Glu Glu Asn Thr Gly Asn Gly Gly Ala Val Thr Ala Asp Asn Pro Lys Asn Glu Asp Glu Val Ala Gln Asn Asp Met Pro Gln 70 65 75 Asn Ala Ala Gly Thr Asp Ser Ser Thr Pro Asn His Thr Pro Asp Pro Asn Met Leu Ala Gly Asn Met Glu Asn Gln Ala Thr Asp Ala Gly Glu 100 105 Ser Ser Gln Pro Ala Asn Gln Pro Asp Met Ala Asn Ala Ala Asp Gly 115 120 Met Gln Gly Asp Asp Pro Ser Ala Gly Gln Asn Ala Gly Asn Thr 135 Ala Ala Gln Gly Ala Asn Gln Ala Gly Asn Asn Gln Ala Ala Gly Ser 150 155 Ser Asp Pro Ile Pro Ala Ser Asn Pro Ala Pro Ala Asn Gly Gly Ser Asn Phe Gly Arg Val Asp Leu Ala Asn Gly Val Leu Ile Asp Gly Pro 185 180

Ser Gln Asn Ile Thr Leu Thr His Cys Lys Gly Asp Ser Cys Ser Gly

205

200

Asn Asn Phe Leu Asp Glu Glu Val Gln Leu Lys Ser Glu Phe Glu Lys 210 215 Leu Ser Asp Ala Asp Lys Ile Ser Asn Tyr Lys Lys Asp Gly Lys Asn 230 225 235 240 Asp Lys Phe Val Gly Leu Val Ala Asp Ser Val Gln Met Lys Gly Ile 245 250 Asn Gln Tyr Ile Ile Phe Tyr Lys Pro Lys Pro Thr Ser Phe Ala Arg 260 265 Phe Arg Arg Ser Ala Arg Ser Arg Arg Ser Leu Pro Ala Glu Met Pro 280 Leu Ile Pro Val Asn Gln Ala Asp Thr Leu Ile Val Asp Gly Glu Ala 295 290 Val Ser Leu Thr Gly His Ser Gly Asn Ile Phe Ala Pro Glu Gly Asn 305 310 315 Tyr Arg Tyr Leu Thr Tyr Gly Ala Glu Lys Leu Pro Gly Gly Ser Tyr 325 330 Ala Leu Arg Val Gln Gly Glu Pro Ala Lys Gly Glu Met Leu Ala Gly 340 345 Ala Ala Val Tyr Asn Gly Glu Val Leu His Phe His Thr Glu Asn Gly 360 Arg Pro Tyr Pro Thr Arg Gly Arg Phe Ala Ala Lys Val Asp Phe Gly 370 375 Ser Lys Ser Val Asp Gly Ile Ile Asp Ser Gly Asp Asp Leu His Met 385 395 Gly Thr Gln Lys Phe Lys Ala Ala Ile Asp Gly Asn Gly Phe Lys Gly 405 410 Thr Trp Thr Glu Asn Gly Ser Gly Asp Val Ser Gly Lys Phe Tyr Gly 420 425 Pro Ala Gly Glu Glu Val Ala Gly Lys Tyr Ser Tyr Arg Pro Thr Asp 440 Ala Glu Lys Gly Gly Phe Gly Val Phe Ala Gly Lys Lys Glu Gln Asp 455 460 <210> <211> 644 <212> PRT <213> N. meningitidis

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Ala Thr Asp Lys Pro Lys Asn Glu Asp Glu Gly Ala Gln Asn Asp Met 65 70 75 80

Pro Gln Asn Ala Ala Asp Thr Asp Ser Leu Thr Pro Asn His Thr Pro 85 90 95

Ala Ser Asn Met Pro Ala Gly Asn Met Glu Asn Gln Ala Pro Asp Ala 100 105 110

Gly Glu Ser Glu Gln Pro Ala Asn Gln Pro Asp Met Ala Asn Thr Ala 115 120 125

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Asn Thr Ala Ala Gln Gly Thr Asn Gln Ala Glu Asn Asn Gln Thr Ala 145 150 155 160

Gly Ser Gln Asn Pro Ala Ser Ser Thr Asn Pro Ser Ala Thr Asn Ser 165 170 175

Gly Gly Asp Phe Gly Arg Thr Asn Val Gly Asn Ser Val Val Ile Asp 180 185 190

Gly Pro Ser Gln Asn Ile Thr Leu Thr His Cys Lys Gly Asp Ser Cys 195 200 205

Ser Gly Asn Asn Phe Leu Asp Glu Glu Val Gln Leu Lys Ser Glu Phe 210 215 220

Glu Lys Leu Ser Asp Ala Asp Lys Ile Ser Asn Tyr Lys Lys Asp Gly
225 230 235 240

Lys Asn Asp Gly Lys Asn Asp Lys Phe Val Gly Leu Val Ala Asp Ser 245 250 255

Val Gln Met Lys Gly Ile Asn Gln Tyr Ile Ile Phe Tyr Lys Pro Lys 260 265 270

Pro Thr Ser Phe Ala Arg Phe Arg Arg Ser Ala Arg S